

IN THE CLAIMS

Please amend Claim 35 as follows.

35. (Currently Amended) A peptide that:

a) is at least 8 amino acid residues long and is a fragment of a mutant protein arising from a frameshift mutation in ~~a gene~~ the TGF- $\beta$ -RII gene or the BAX gene of a cancer cell having a protein sequence that consists of a mutant part and a normal part;

31 b) includes at least one amino acid residue of the mutant part of the protein sequence;

c) comprises 0-10 amino acid residues from the carboxyl terminus of the normal part of the protein sequence preceding the amino terminus of the mutant part of the protein sequence and may further extend to the carboxyl terminus of the mutant part of the protein sequence, as determined by a new stop codon generated by the frameshift mutation; and

d) induces T cell responses, either in its full length form or after processing by an antigen presenting cell;

wherein the mutant part of the protein sequence has a sequence chosen from sequence identity nos. 1-459.

Claims 36-73 (Original).